

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/560, 470
Source: IFWP
Date Processed by STIC: 12/28/2005

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 12/28/2005

PATENT APPLICATION: US/10/560,470

TIME: 12:42:59

Input Set : A:\Lannigan.00910-05.txt

Output Set: N:\CRF4\12282005\J560470.raw

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3 <110> APPLICANT: University of Virginia Patent Foundation
4     Lannigan-Macara, Deborah A.
5     Henrich, Lorin M.
6     Smith, Jeffrey A.
8 <120> TITLE OF INVENTION: ERK7 and ERK8, Novel Diagnostic Markers for Cancer
10 <130> FILE REFERENCE: 00910-05
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/560,470
C--> 12 <141> CURRENT FILING DATE: 2005-12-14
12 <150> PRIOR APPLICATION NUMBER: US 60478,992
13 <151> PRIOR FILING DATE: 2003-06-17
15 <150> PRIOR APPLICATION NUMBER: PCT/US2004/019181
16 <151> PRIOR FILING DATE: 2004-06-17
18 <160> NUMBER OF SEQ ID NOS: 6
20 <170> SOFTWARE: PatentIn version 3.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 484
24 <212> TYPE: PRT
25 <213> ORGANISM: Homo sapiens
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34          20          25          30
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38          35          40          45
41 Arg Asp Lys Thr Asp Ala Gln Arg Thr Phe Arg Glu Ile Thr Leu Leu
42          50          55          60
45 Gln Glu Phe Gly Asp His Pro Asn Ile Ile Ser Leu Leu Asp Val Ile
46 65          70          75          80
49 Arg Ala Glu Asn Asp Arg Asp Ile Tyr Leu Val Phe Glu Phe Met Asp
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53 Thr Asp Leu Asn Ala Val Ile Arg Lys Gly Gly Leu Leu Gln Asp Val
54          100         105         110
57 His Val Arg Ser Ile Phe Tyr Gln Leu Leu Arg Ala Thr Arg Phe Leu
58          115         120         125
61 His Ser Gly His Val Val His Arg Asp Gln Lys Pro Ser Asn Val Leu
62          130         135         140
65 Leu Asp Ala Asn Cys Thr Val Lys Leu Cys Asp Phe Gly Leu Ala Arg
66 145         150         155         160
69 Ser Leu Gly Asp Leu Pro Glu Gly Pro Glu Asp Gln Ala Val Thr Glu
70          165         170         175
73 Tyr Val Ala Glu Asp Leu Leu Ala Leu Gly Ser Gly Cys Arg Ala Ser
74          180         185         190

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77 Val Leu His Gln Leu Gly Ser Arg Pro Arg Gln Thr Leu Asp Ala Leu
78      195      200      205
81 Leu Pro Pro Asp Thr Ser Pro Glu Ala Leu Asp Leu Leu Arg Arg Leu
82      210      215      220
85 Leu Val Phe Ala Pro Asp Lys Arg Leu Ser Ala Thr Gln Ala Leu Gln
86 225      230      235      240
89 His Pro Tyr Val Gln Arg Phe His Cys Pro Ser Asp Glu Trp Ala Arg
90      245      250      255
93 Glu Ala Asp Val Arg Pro Arg Ala His Glu Gly Val Gln Leu Ser Val
94      260      265      270
97 Pro Glu Tyr Arg Ser Arg Val Tyr Gln Met Ile Leu Glu Cys Gly Gly
98      275      280      285
101 Ser Ser Gly Thr Ser Arg Glu Lys Gly Pro Glu Gly Val Ser Pro Ser
102      290      295      300
105 Gln Ala His Leu His Lys Pro Arg Ala Asp Pro Gln Leu Pro Ser Arg
106 305      310      315      320
109 Thr Pro Val Gln Gly Pro Arg Pro Arg Pro Gln Ser Ser Pro Gly His
110      325      330      335
113 Asp Pro Ala Glu His Glu Ser Pro Arg Ala Ala Lys Asn Val Pro Arg
114      340      345      350
117 Gln Asn Ser Ala Pro Leu Leu Gln Thr Ala Leu Leu Gly Asn Gly Glu
118      355      360      365
121 Arg Pro Pro Gly Ala Lys Glu Ala Pro Pro Leu Thr Leu Ser Leu Val
122      370      375      380
125 Lys Pro Ser Gly Arg Gly Ala Ala Pro Ser Leu Thr Ser Gln Ala Ala
126 385      390      395      400
129 Ala Gln Val Ala Asn Gln Ala Leu Ile Arg Gly Asp Trp Asn Arg Gly
130      405      410      415
133 Gly Gly Val Arg Val Ala Ser Val Gln Gln Val Pro Pro Arg Leu Pro
134      420      425      430
137 Pro Glu Ala Arg Pro Gly Arg Arg Met Phe Ser Thr Ser Ala Leu Gln
138      435      440      445
141 Gly Ala Gln Gly Gly Ala Arg Ala Leu Leu Gly Gly Tyr Ser Gln Ala
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168 Asp Arg Arg Thr Gly Glu Val Val Ala Ile Lys Lys Ile Phe Asp Ala
169      35      40      45
172 Phe Arg Asp Gln Thr Asp Ala Gln Arg Thr Phe Arg Glu Ile Met Leu

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173      50      55      60
176 Leu Arg Glu Phe Gly Gly His Pro Asn Ile Ile Arg Leu Leu Asp Val
177 65      70      75      80
180 Ile Pro Ala Lys Asn Asp Arg Asp Ile Tyr Leu Val Phe Glu Ser Met
181      85      90      95
184 Asp Thr Asp Leu Asn Ala Val Ile Gln Lys Gly Arg Leu Leu Glu Asp
185      100      105      110
188 Ile His Lys Arg Cys Ile Phe Tyr Gln Leu Leu Arg Ala Thr Lys Phe
189      115      120      125
192 Ile His Ser Gly Arg Val Ile His Arg Asp Gln Lys Pro Ala Asn Val
193      130      135      140
196 Leu Leu Asp Ala Ala Cys Arg Val Lys Leu Cys Asp Phe Gly Leu Ala
197 145      150      155      160
200 Arg Ser Leu Ser Asp Phe Pro Glu Gly Pro Gly Gly Gln Ala Leu Thr
201      165      170      175
204 Glu Tyr Val Ala Thr Arg Trp Tyr Arg Ala Pro Glu Val Leu Leu Ser
205      180      185      190
208 Ser Arg Trp Tyr Thr Pro Gly Val Asp Met Trp Ser Leu Gly Cys Ile
209      195      200      205
212 Leu Gly Glu Met Leu Arg Gly Gln Pro Leu Phe Pro Gly Thr Ser Thr
213      210      215      220
216 Phe His Gln Leu Glu Leu Ile Leu Glu Thr Ile Pro Leu Pro Ser Met
217 225      230      235      240
220 Glu Glu Leu Gln Gly Leu Gly Ser Asp Tyr Ser Ala Leu Ile Leu Gln
221      245      250      255
224 Asn Leu Gly Ser Arg Pro Arg Gln Thr Leu Asp Ala Leu Leu Pro Pro
225      260      265      270
228 Asp Thr Pro Pro Glu Ala Leu Asp Leu Leu Lys Arg Leu Leu Ala Phe
229      275      280      285
232 Ala Pro Asp Lys Arg Leu Ser Ala Glu Gln Ala Leu Gln His Pro Tyr
233      290      295      300
236 Val Gln Arg Phe His Cys Pro Asp Arg Glu Trp Thr Arg Gly Ser Asp
237 305      310      315      320
240 Val Arg Leu Pro Val His Glu Gly Asp Gln Leu Ser Ala Pro Glu Tyr
241      325      330      335
244 Arg Asn Arg Leu Tyr Gln Met Ile Leu Glu Arg Arg Arg Asn Ser Arg
245      340      345      350
248 Ser Pro Arg Glu Glu Asp Leu Gly Val Val Ala Ser Arg Ala Glu Leu
249      355      360      365
252 Arg Ala Ser Gln Arg Gln Ser Leu Lys Pro Gly Val Leu Pro Gln Val
253      370      375      380
256 Leu Ala Glu Thr Pro Ala Arg Lys Arg Gly Pro Lys Pro Gln Asn Gly
257 385      390      395      400
260 His Gly His Asp Pro Glu His Val Glu Val Arg Arg Gln Ser Ser Asp
261      405      410      415
264 Pro Leu Tyr Gln Leu Pro Pro Pro Gly Ser Gly Glu Arg Pro Pro Gly
265      420      425      430
268 Ala Thr Gly Glu Pro Pro Ser Ala Pro Ser Gly Val Lys Thr His Val
269      435      440      445

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272 Arg Ala Val Ala Pro Ser Leu Thr Ser Gln Ala Ala Ala Gln Ala Ala
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276 Asn Gln Pro Leu Ile Arg Ser Asp Pro Ala Arg Gly Gly Gly Pro Arg
277 465                      470                      475                      480
280 Ala Val Gly Ala Arg Arg Val Pro Ser Arg Leu Pro Arg Glu Ala Pro
281      485                      490                      495
284 Glu Pro Arg Pro Gly Arg Arg Met Phe Gly Ile Ser Val Ser Gln Gly
285      500                      505                      510
288 Ala Gln Gly Ala Ala Arg Ala Ala Leu Gly Gly Tyr Ser Gln Ala Tyr
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315 Asp Arg Arg Thr Gly Glu Val Val Ala Ile Lys Lys Ile Phe Asp Ala
316      35      40      45
319 Phe Arg Asp Gln Ile Asp Ala Gln Arg Thr Phe Arg Glu Ile Met Leu
320      50      55      60
323 Leu Lys Glu Phe Gly Gly His Pro Asn Ile Ile Arg Leu Leu Asp Val
324 65      70      75      80
327 Ile Pro Ala Lys Asn Asp Arg Asp Ile Tyr Leu Val Phe Glu Ser Met
328      85      90      95
331 Asp Thr Asp Leu Asn Ala Val Ile Gln Lys Gly Arg Leu Leu Lys Asp
332      100     105     110
335 Ile His Lys Arg Cys Ile Phe Tyr Gln Leu Leu Arg Ala Thr Lys Phe
336      115     120     125
339 Ile His Ser Gly Arg Val Ile His Arg Asp Gln Lys Pro Ala Asn Val
340      130     135     140
343 Leu Leu Asp Ser Ala Cys Arg Val Lys Leu Cys Asp Phe Gly Leu Ala
344 145     150     155     160
347 Arg Ser Leu Gly Asp Leu Pro Glu Gly Pro Gly Gly Gln Ala Leu Thr
348      165     170     175
351 Glu Tyr Val Ala Thr Arg Trp Tyr Arg Ala Pro Glu Val Leu Leu Ser
352      180     185     190
355 Ser Arg Trp Tyr Thr Pro Gly Val Asp Met Trp Ser Leu Gly Cys Ile
356      195     200     205
359 Leu Gly Glu Met Leu Arg Gly Gln Pro Leu Phe Pro Gly Thr Ser Thr
360      210     215     220
363 Phe His Gln Leu Glu Leu Ile Leu Lys Thr Ile Pro Leu Pro Ser Met
364 225     230     235     240

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367 Glu Glu Leu Gln Asp Leu Gly Ser Asp Tyr Ser Ala Leu Ile Leu Gln
368                245                250                255
371 Asn Leu Gly Ser Arg Pro Gln Gln Thr Leu Asp Ala Leu Leu Pro Pro
372                260                265                270
375 Asp Thr Pro Pro Glu Ala Leu Asp Leu Leu Lys Arg Leu Leu Ala Phe
376                275                280                285
379 Ala Pro Asp Lys Arg Leu Ser Ala Glu Gln Ala Leu Gln His Pro Tyr
380                290                295                300
383 Val Gln Arg Phe His Cys Pro Asp Arg Glu Trp Ala Arg Glu Ser Asp
384 305                310                315                320
387 Val Arg Leu Pro Val His Glu Gly Asp Gln Leu Ser Ala Pro Glu Tyr
388                325                330                335
391 Arg Lys Arg Leu Tyr Gln Ile Ile Leu Glu Gln Ser Gly Asn Ser Arg
392                340                345                350
395 Ser Pro Arg Glu Glu Gly Leu Gly Val Val Ala Ser Arg Ala Glu Leu
396                355                360                365
399 Arg Ala Ser Pro Ala Arg Thr Gln Ser Leu Lys Ser Gly Val Leu Pro
400                370                375                380
403 Gln Val Pro Ala Glu Thr Pro Ala Arg Lys Arg Gly Pro Lys Pro Pro
404 385                390                395                400
407 Arg Ser Pro Gly His Asp Pro Glu His Val Glu Val Arg Arg Gln Ser
408                405                410                415
411 Ser Asp Pro Leu Phe Gln Leu Pro Pro Pro Gly Arg Gly Glu Arg Pro
412                420                425                430
415 Pro Gly Ala Thr Gly Gln Pro Pro Ser Ala Pro Ser Gly Val Lys Thr
416                435                440                445
419 Gln Val Arg Ala Met Ala Pro Ser Leu Thr Ser Gln Ala Glu Ala Gln
420                450                455                460
423 Ala Ala Asn Gln Ala Leu Ile Arg Ser Asp Pro Ala Arg Gly Gly Gly
424 465                470                475                480
427 Pro Arg Ala Val Gly Ala Arg Arg Val Pro Ser Arg Leu Pro Arg Glu
428                485                490                495
431 Ala Pro Glu Pro Arg Pro Gly Arg Arg Met Phe Gly Ile Ser Val Ser
432                500                505                510
435 Gln Gly Ala Gln Gly Ala Ala Arg Ala Ala Leu Gly Gly Tyr Ser Gln
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VERIFICATION SUMMARY

PATENT APPLICATION: **US/10/560,470**

DATE: 12/28/2005

TIME: 12:43:00

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date